Figure S4

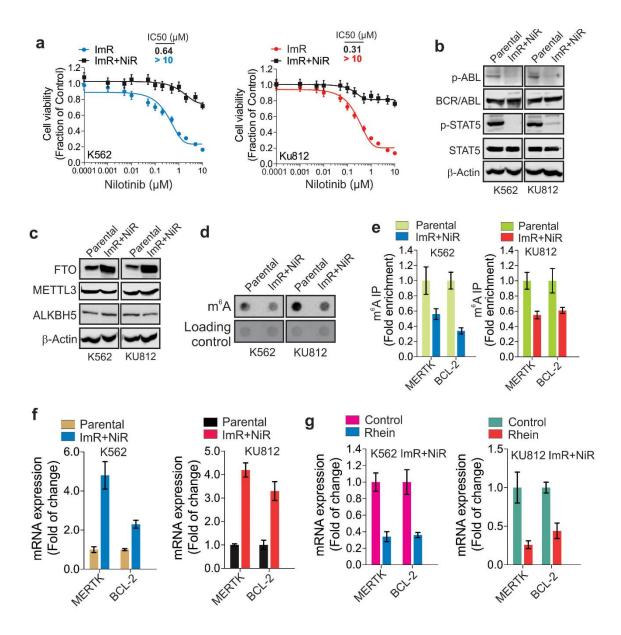


Figure S4. FTO upregulation and m⁶A demethylation occur in second-line resistant cells. **a** CCK-8 assays in first- (ImR) and second- (ImR+NiR) line K562 and KU812 resistant cells treated with nilotinib for 72 hours. The data represent two independent experiments with 8 repeats in total. **b,c** Western blotting of parental and ImR+NiR cells. **d** Dotblotting of parental and ImR+NiR cells. **e** The mRNA from parental and ImR+NiR cells was subjected to immunoprecipitation (IP) by antim⁶A. The eluted RNA was subjected to cDNA synthesis followed by qPCR for gene expression.

f,**g** qPCR of parental vs ImR+NiR cells (**f**) or ImR+NiR cells treated with 25 μ M rhein for 48 hours (**g**).

In \mathbf{b} - \mathbf{g} , data represent three independent experiments.